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Bacterial Cytological Profiling Reveals the Mechanism of Action of Anticancer Metal Complexes

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Calculation of the ratio of DNA and RNA nucleotides to metal based on data reported by DeRose *et al.*¹:

The platinum content measured after 12 hours treatment of 100 μM cisplatin was used for this calculation.

$$\text{Pt} / \text{DNA nt} = 60.2 \times 10^{-5}$$

$$\text{DNA nt} / \text{Pt} = \frac{1}{60.2 \times 10^{-5}} = 1661:1$$

$$\text{Pt} / \text{RNA nt} = 15.7 \times 10^{-5}$$

$$\text{RNA nt} / \text{Pt} = \frac{1}{15.7 \times 10^{-5}} = 6369:1$$

Calculation of the ratio of DNA and RNA nucleotides to metal based on data reported from an IVTT assay² (at the IC_{50}):

$$\text{DNA bp} / \text{Ru} = 570:1$$

$$\text{DNA nt} / \text{Ru} = 1140:1$$

$$\text{DNA bp} / \text{Pt} = 300:1$$

$$\text{DNA nt} / \text{Pt} = 600:1$$

$$\text{RNA nt} / \text{Ru} = 610:1$$

$$\text{RNA nt} / \text{Pt} = 820:1$$

Calculation of the ratio of DNA and RNA nucleotides to metal at the IC_{50} to Dendra2 production inhibition based on data collected in AAS:

The DNA and RNA nt / mc ratio collected by AAS at 20 μM is as follows:

$$\text{DNA nt} / \text{Ru} = 2000:1$$

$$\text{DNA nt} / \text{Pt} = 3000:1$$

$$\text{RNA nt} / \text{Ru} = 3800:1$$

$$\text{RNA nt} / \text{Pt} = 4700:1$$

Assuming the uptake is linearly proportional to the dosed concentration, the IC_{50} value to inhibit Dendra2 production in *E. coli* is 77 μM for compound **1** and 85 μM for cisplatin. Thus, a correction coefficient can be calculated to adjust the metal content to what would be obtained at the higher compound dose:

$$\text{Compound 1: } \frac{77}{20} = 3.8$$

$$\text{Cisplatin: } \frac{85}{20} = 4.3$$

Then, at the IC_{50} to inhibit Dendra2 production, the theoretical metal to DNA or RNA nucleotides to metal ratio can be calculated as follows:

$$\text{DNA nt / Ru} = \frac{2000}{3.8} = 520 : 1$$

$$\text{RNA nt / Ru} = \frac{3800}{3.8} = 1000 : 1$$

$$\text{DNA nt / Pt} = \frac{3000}{4.3} = 700 : 1$$

$$\text{RNA nt / Pt} = \frac{4700}{4.3} = 1090 : 1$$

The average molecular weight values used in calculation for DNA base pairs and RNA nucleotides are 665 g/mol and 340 g/mol.

Table S1. Cellular metal content with different nucleic acids measured by AAS.

	<i>E. coli</i>		HL60	
	Percentage of metal bound with genomic DNA ^a	Percentage of metal bound with total RNA ^b	Percentage of metal bound with genomic DNA ^a	Percentage of metal bound with total RNA ^b
1 light	1.3% ± 0.1%	0.5% ± 0.1%	1.3% ± 0.1%	2.0% ± 0.3%
1 dark	^c	^c	^c	^c
cisplatin	1.0% ± 0.1%	0.7% ± 0.1%	1.1% ± 0.1%	1.5% ± 0.2%

^{a, b}The percentage of metal bound with genomic DNA or total RNA was calculated as follows:

$$\text{Percentage of metal bound with DNA (RNA)} = \frac{\text{Metal measured in DNA (RNA) sample } (\mu\text{mol})}{\text{Metal in DNA sample} + \text{cell sample } (\mu\text{mol})}$$

^c Ruthenium level in DNA and RNA samples were under the detection limit (<2ppb).

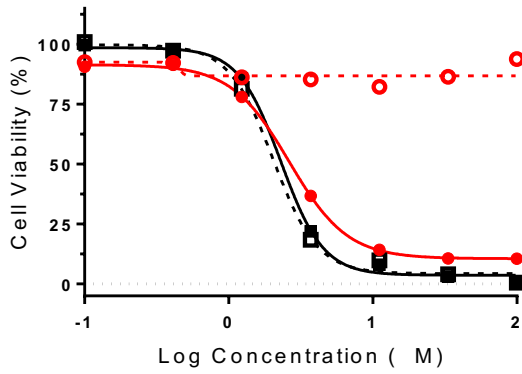


Figure S1. Cytotoxicity dose response of complex **1** (red lines) and cisplatin (black lines) in *E. coli* following photoirradiation (solid lines, filled circles) and dark condition (dashed lines, open circles).

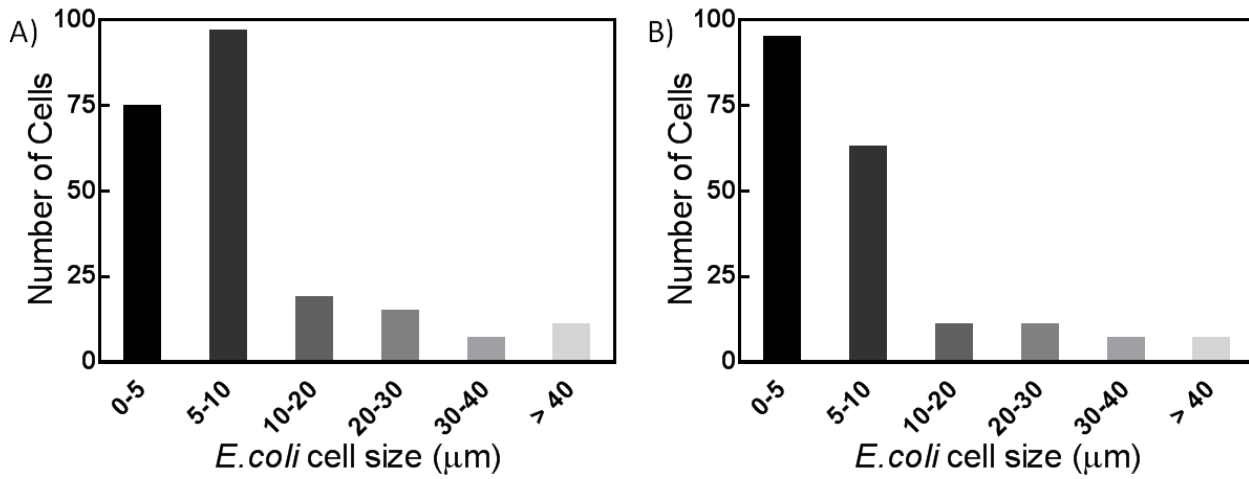


Figure S2. Size distribution histograms of *E. coli* filaments with treatment of A) rifampicin; and B) tetracycline. *E. coli* cells were treated with 3 μM of rifampicin and 48 μM of tetracycline (10x IC₅₀) for 6 hours.

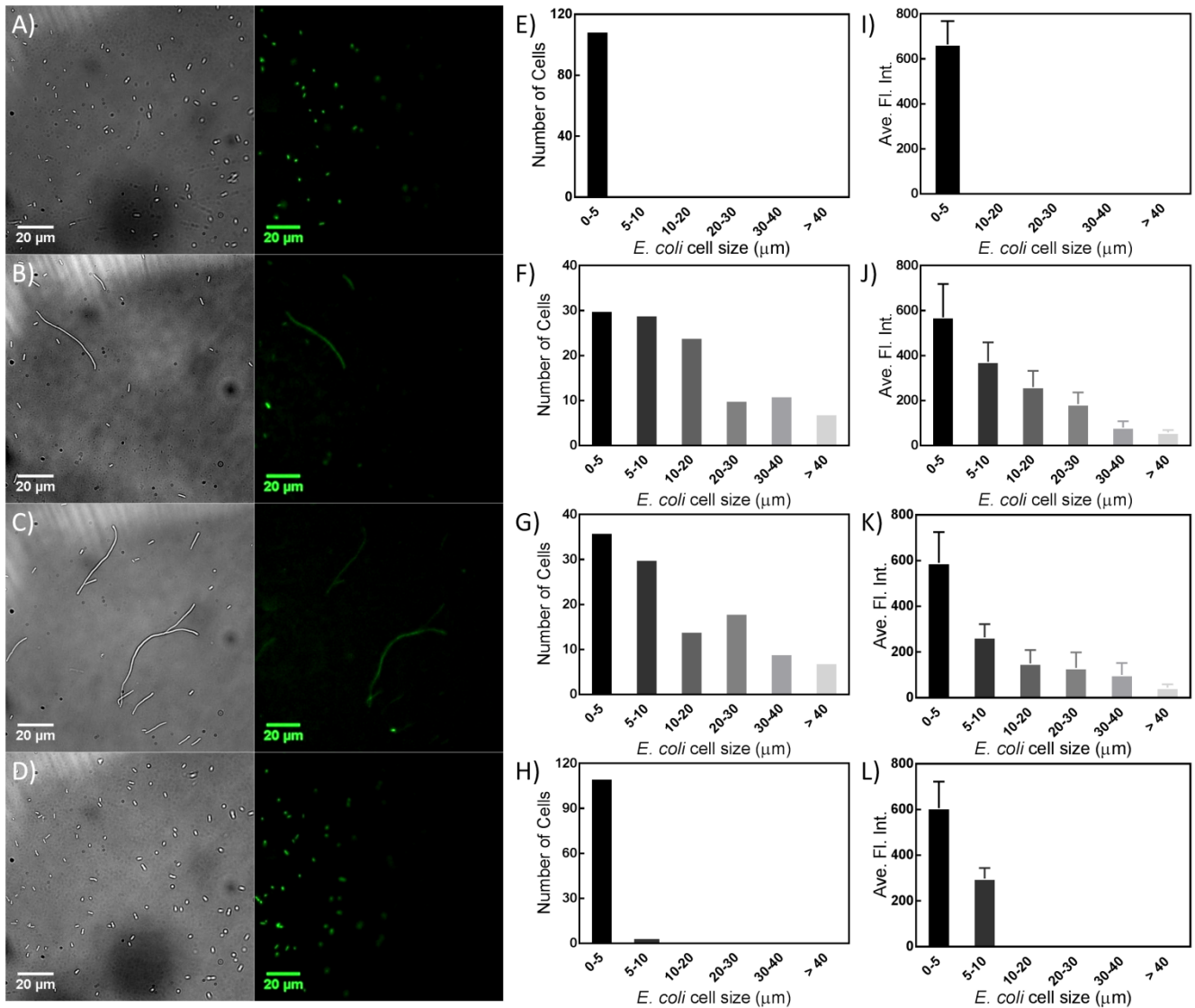


Figure S3. Complex 1 induces filamentous growth and decreased protein production in *E. coli*. A) - D) Bright field and fluorescent imaging of *E. coli* cells under different conditions; E) - H) Size distribution histograms of *E. coli* cells under different treatments; I) - L) Histograms of average fluorescence intensity (Ave. Fl. Int.) correlating to cell size with different treatments. Top through bottom panels: N. C. control, cisplatin, compound 1 with light, and compound 1 without light. Cells were treated with 40 μM each compound for 6 hours before imaging.

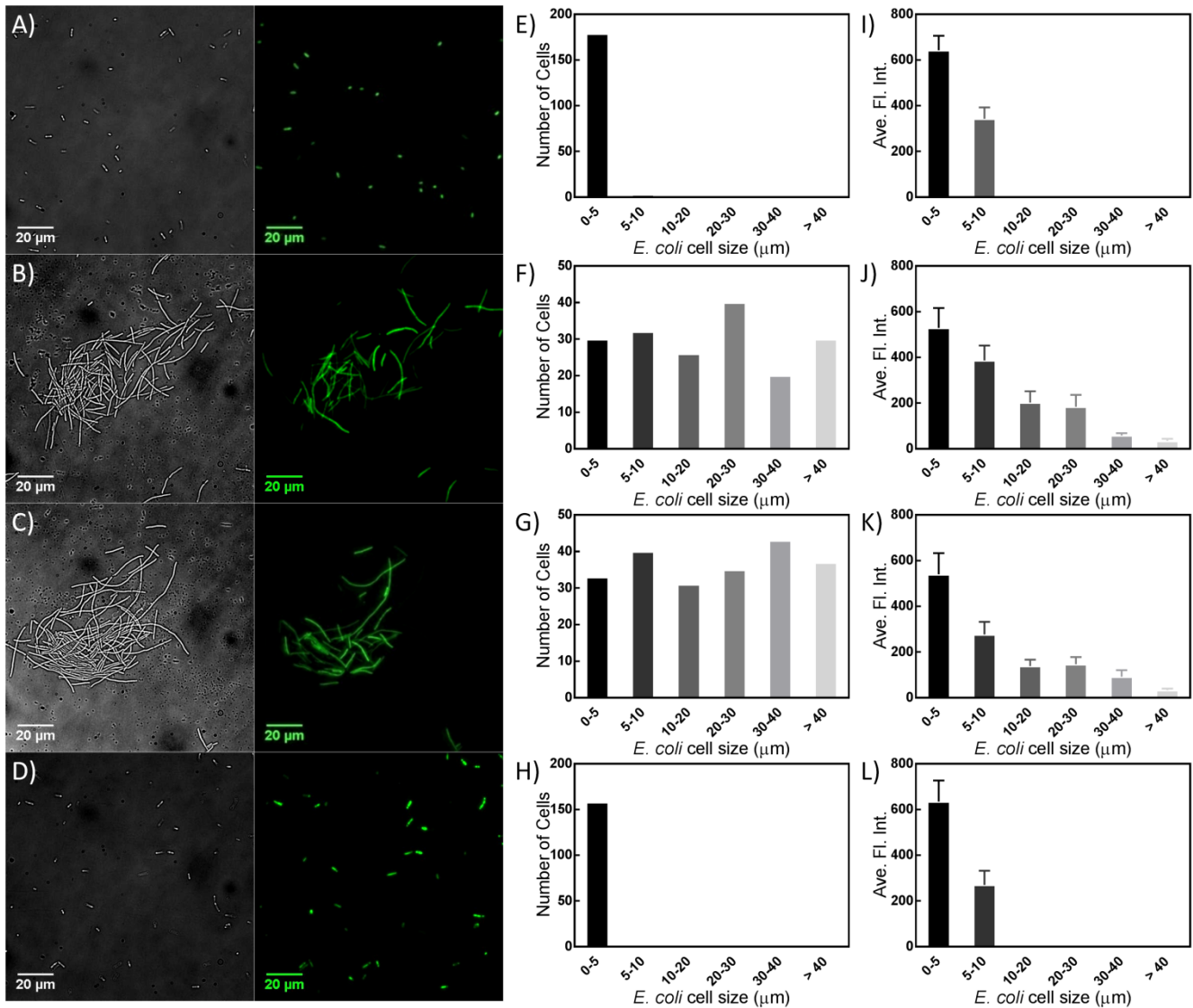


Figure S4. Complex 1 induces filamentous growth and decreased protein production in *E. coli*. A) - D) Bright field and fluorescent imaging of *E. coli* cells under different conditions; E) - H) Size distribution histograms of *E. coli* cells under different treatments; I) - L) Histograms of average fluorescence intensity (Ave. Fl. Int.) correlating to cell size with different treatments. Top through bottom panels: N. C. control, cisplatin, compound 1 with light, and compound 1 without light. Cells were treated with 40 μM each compound for 16 hours before imaging.

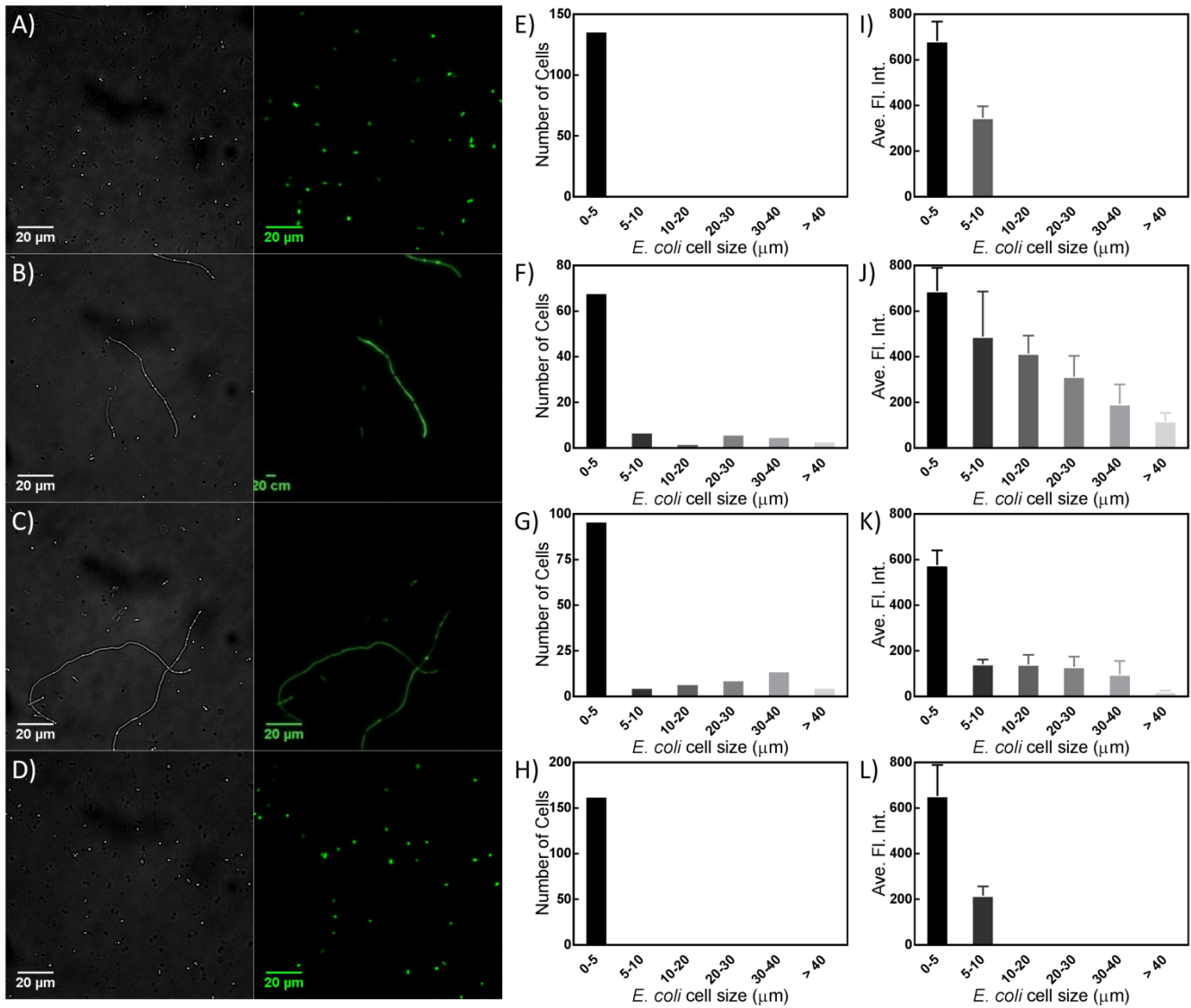


Figure S5. Complex 1 induces filamentous growth and decreased protein production in *E. coli*. A) - D) Bright field and fluorescent imaging of *E. coli* cells under different conditions; E) - H) Size distribution histograms of *E. coli* cells under different treatments; I) - L) Histograms of average fluorescence intensity (Ave. Fl. Int.) correlating to cell size with different treatments. Top through bottom panels: N. C. control, cisplatin, compound 1 with light, and compound 1 without light. Cells were treated with 100 μM each compound for 16 hours before imaging.

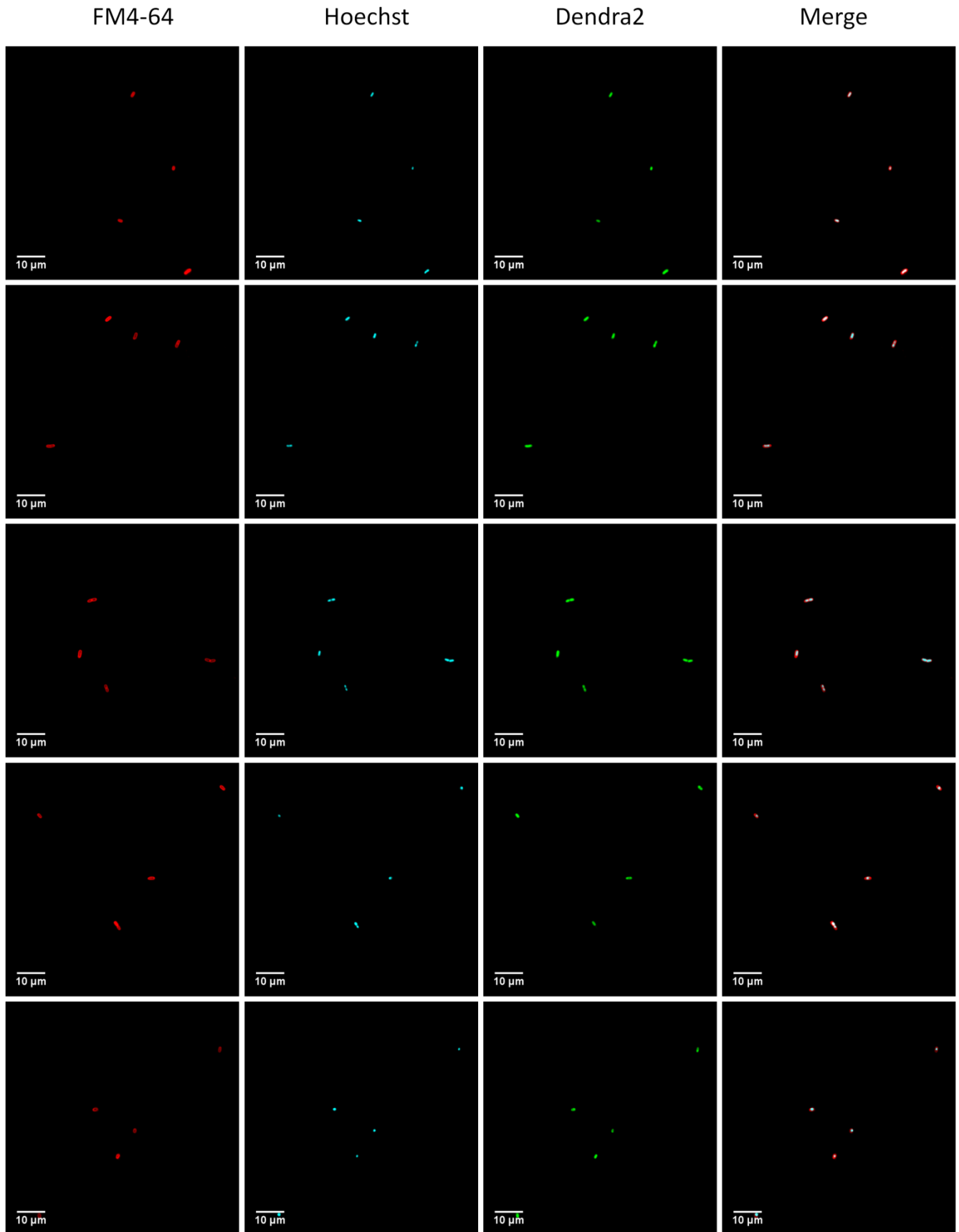


Figure S6. Supplemental fluorescent imaging of *N. C.* control. The merge is the combination of the Hoechst and FM4-64 membrane stain emission data.

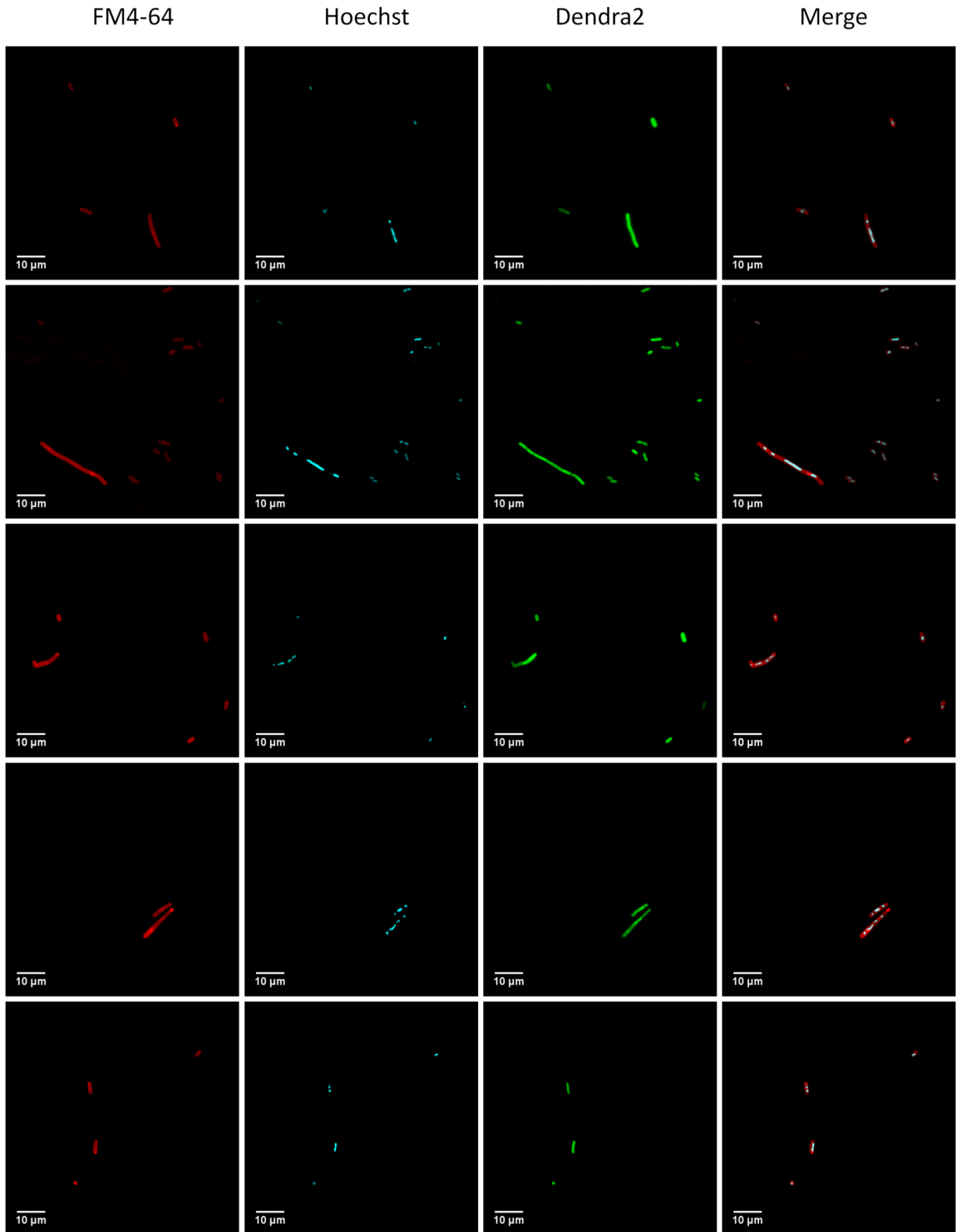


Figure S7. Supplemental fluorescent imaging of cisplatin. Cells were treated MIC for 6 hours before imaging. The merge is the combination of the Hoechst and FM4-64 membrane stain emission data.

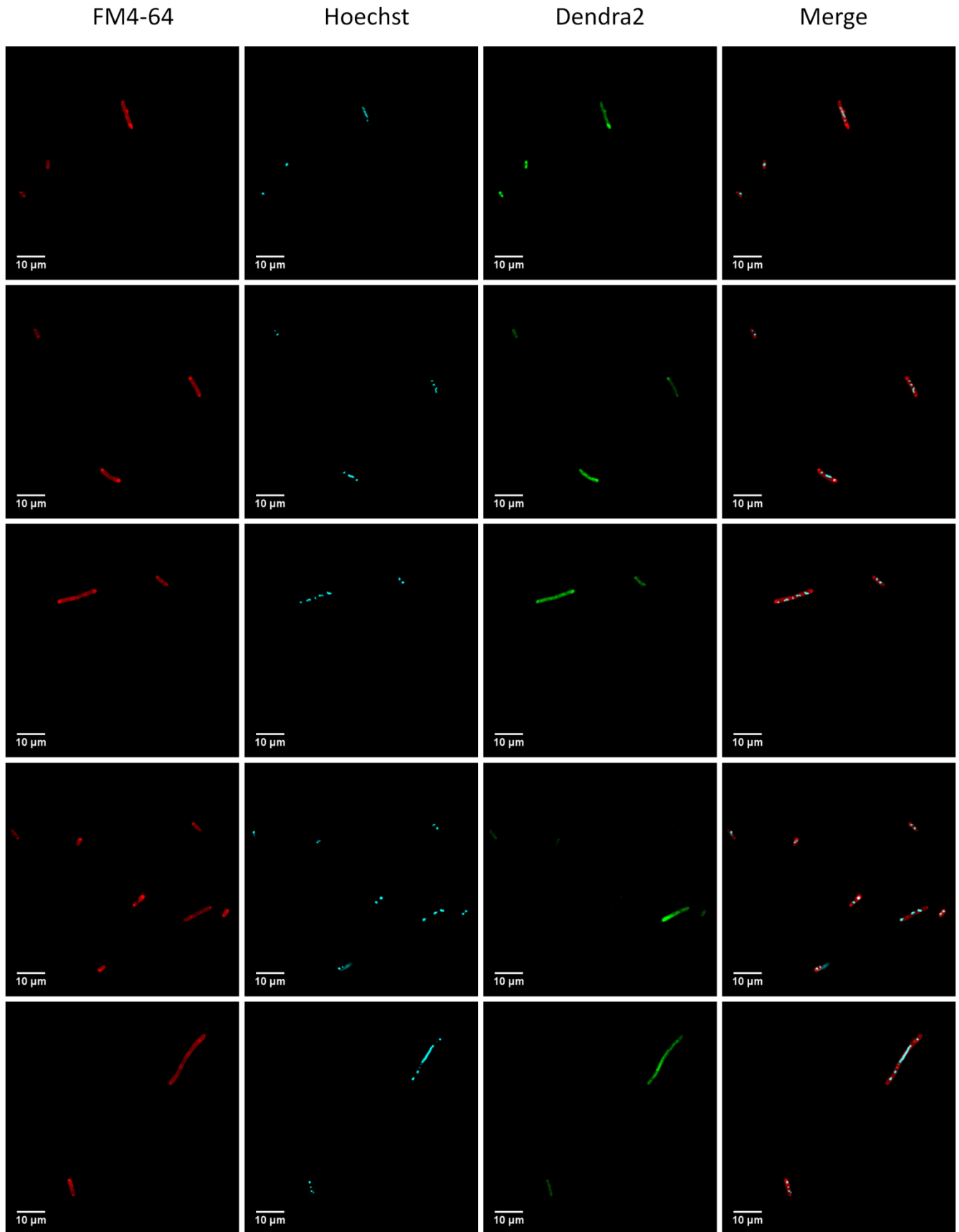


Figure S8. Supplemental fluorescent imaging of compound **1** with light. Cells were treated MIC for 6 hours before imaging. The merge is the combination of the Hoechst and FM4-64 membrane stain emission data.

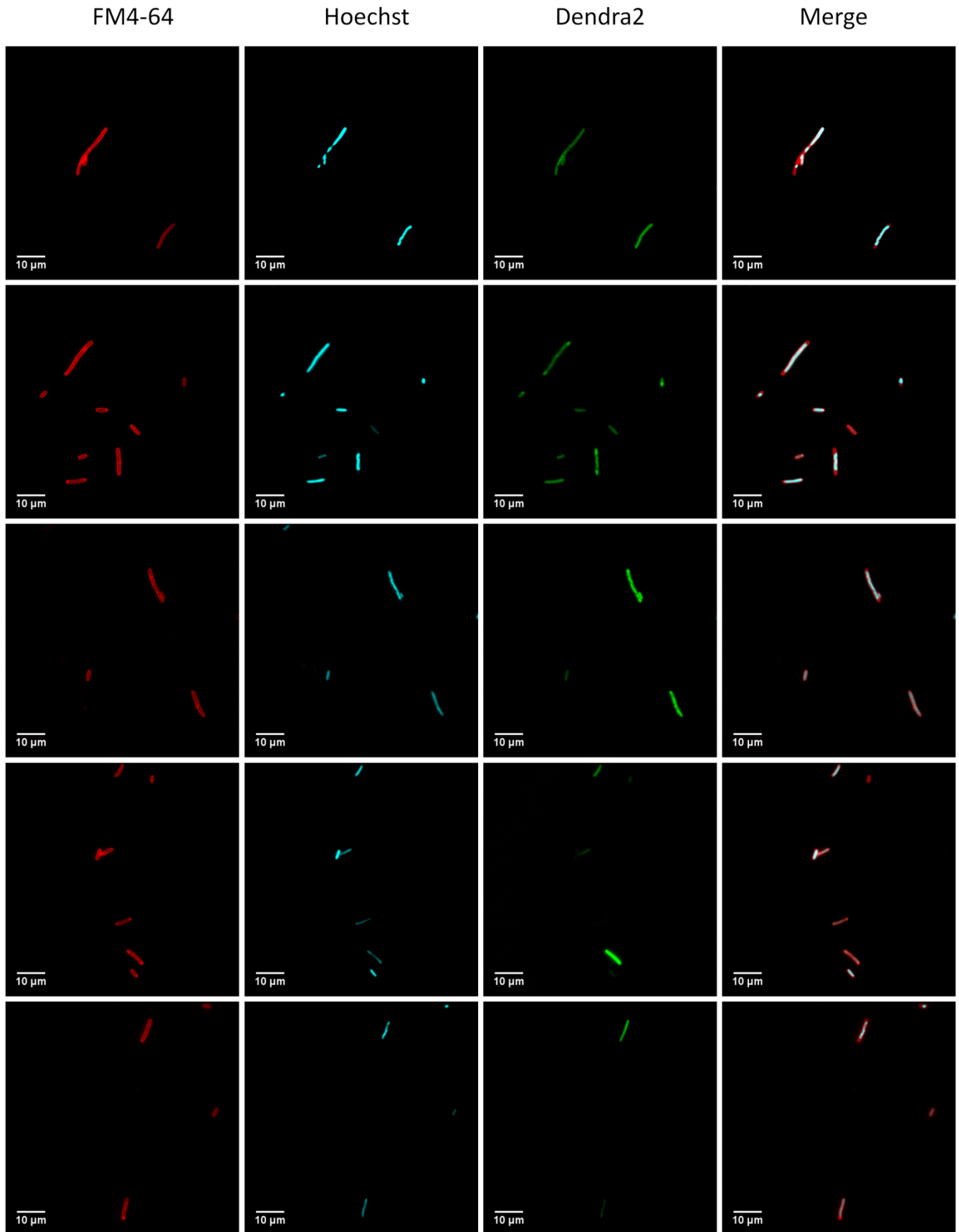


Figure S9. Supplemental fluorescent imaging of rifampicin. Cells were treated 10x MIC for 6 hours before imaging. The merge is the combination of the Hoechst and FM4-64 membrane stain emission data.

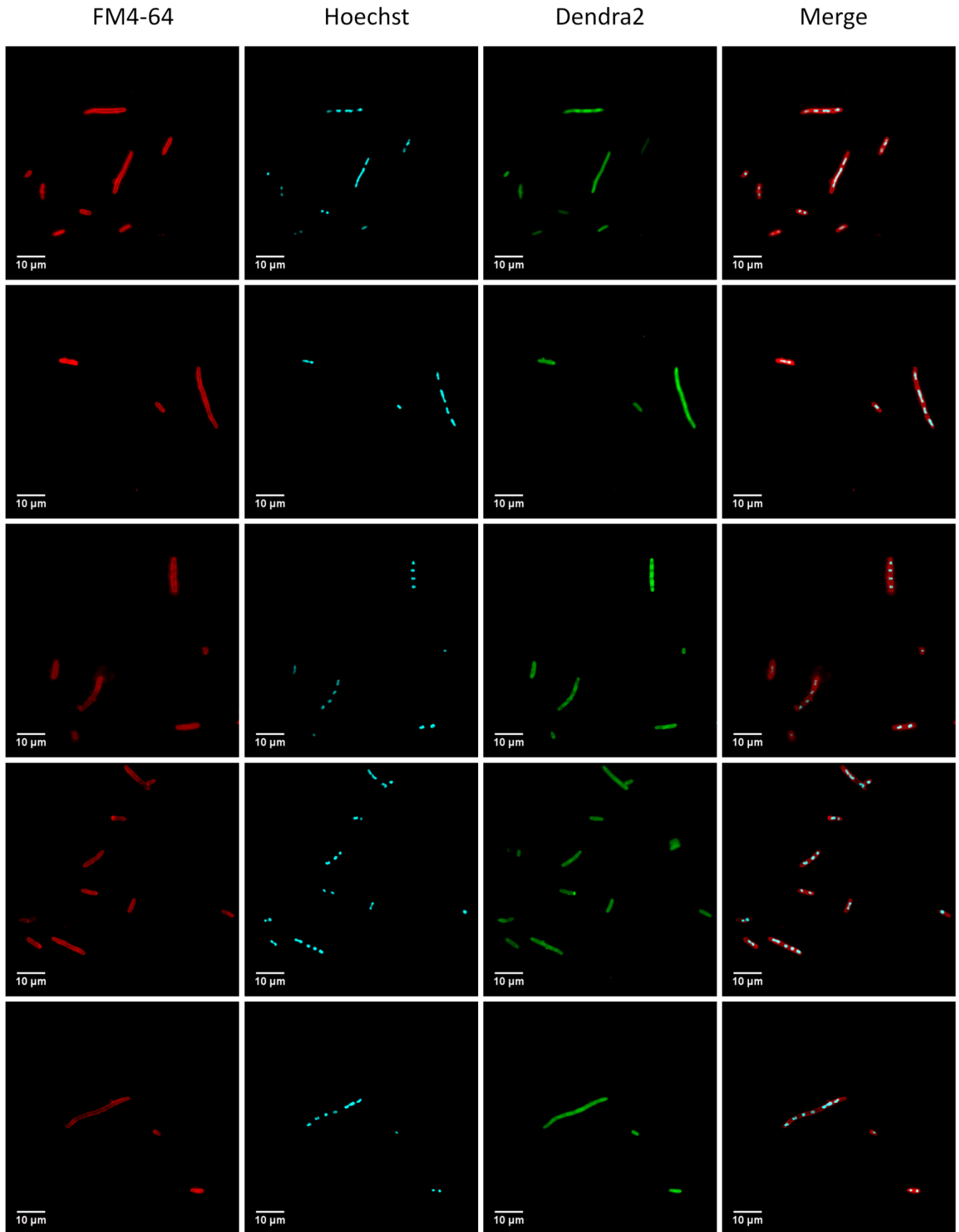


Figure S10. Supplemental fluorescent imaging of tetracycline. Cells were treated MIC for 6 hours before imaging. The merge is the combination of the Hoechst and FM4-64 membrane stain emission data.

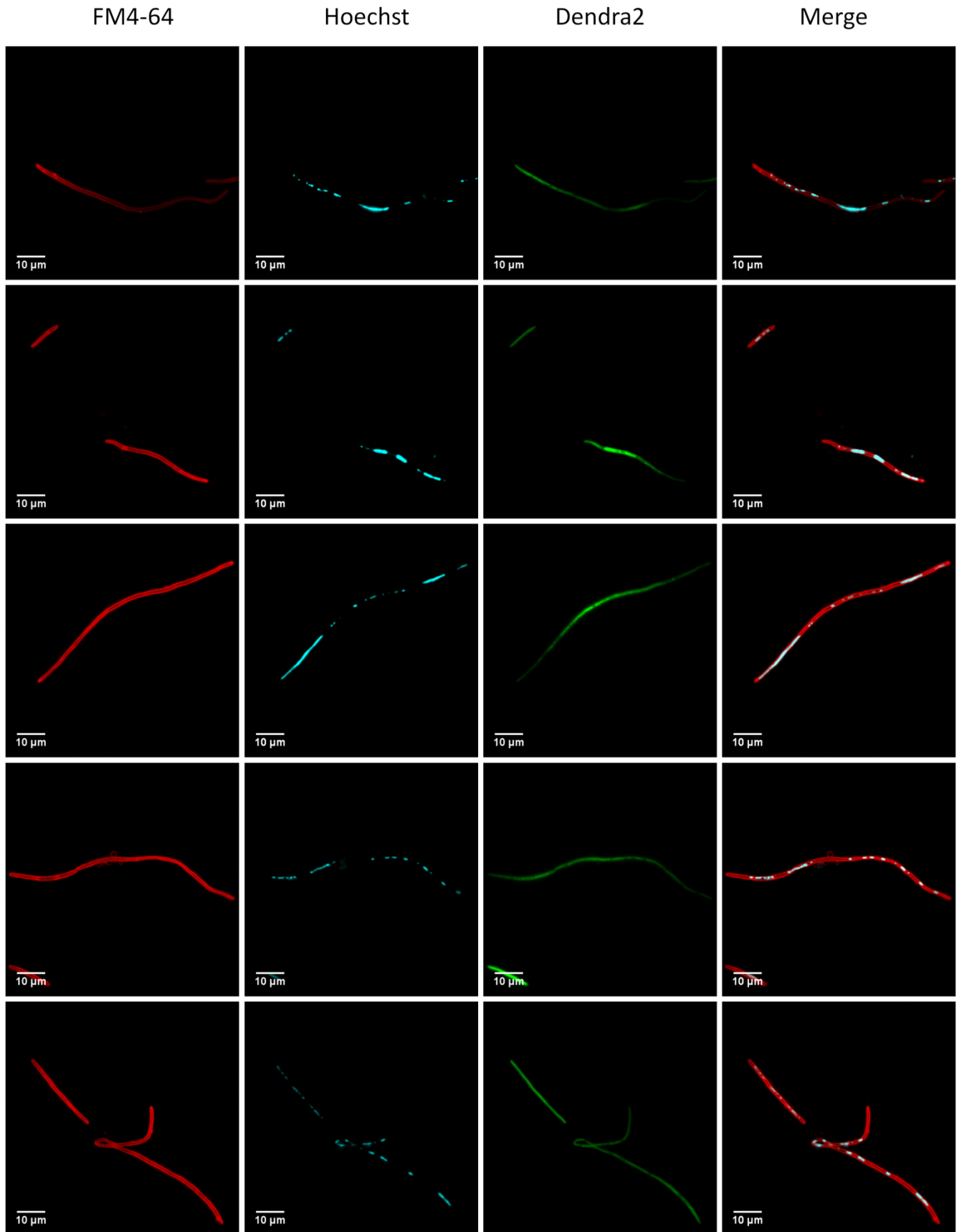


Figure S11. Supplemental fluorescent imaging of nalidixic acid. Cells were treated MIC for 6 hours before imaging. The merge is the combination of the Hoechst and FM4-64 membrane stain emission data.

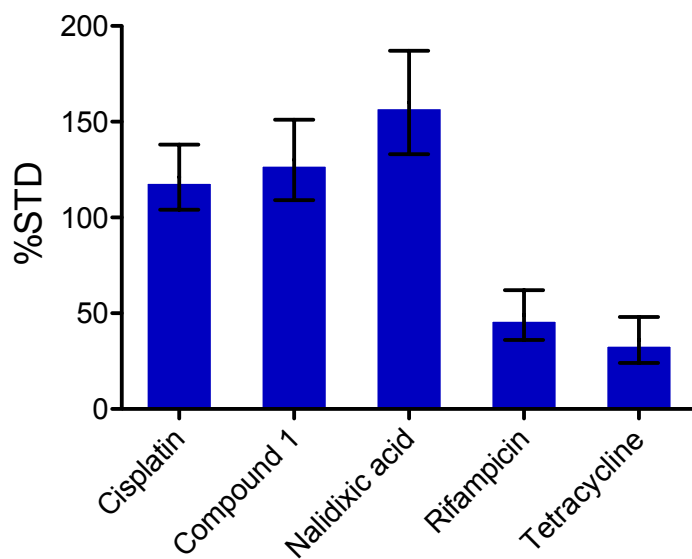


Figure S12. Standard deviations in DNA fragment size for the various compound treatments.

The %STD of DNA fragments was defined and calculated as follows:

$$\%STD \text{ (DNA pieces size)} = \frac{\text{Standard deviation of sizes of DNA pieces}}{\text{Average of sizes of DNA pieces}}$$

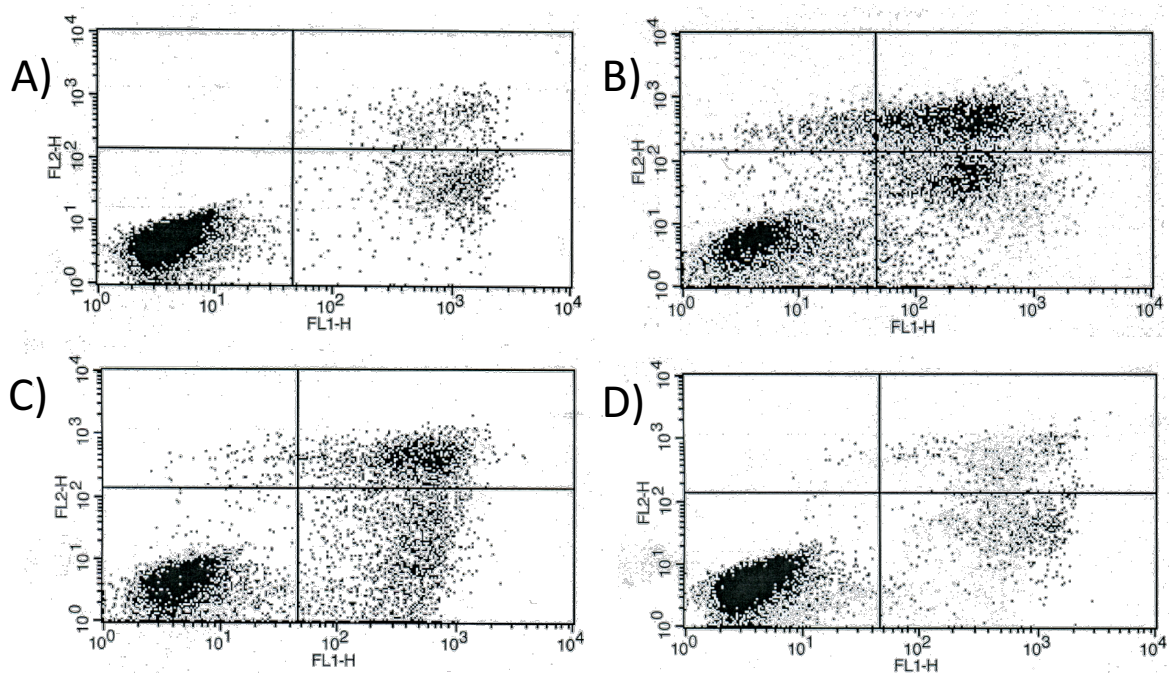


Figure S13. Flow cytometry analysis by FITC/Annexin-V and PI of apoptosis in HL60 cells. FITC-Annexin V (FL1-H) was used in combination with propidium iodide (FL2-H). A) N. C. control; B) cisplatin; C) compound **1** with light irradiation; D) compound **1** in dark. HL60 cells were treated with 20 μ M compounds for 24 hours.

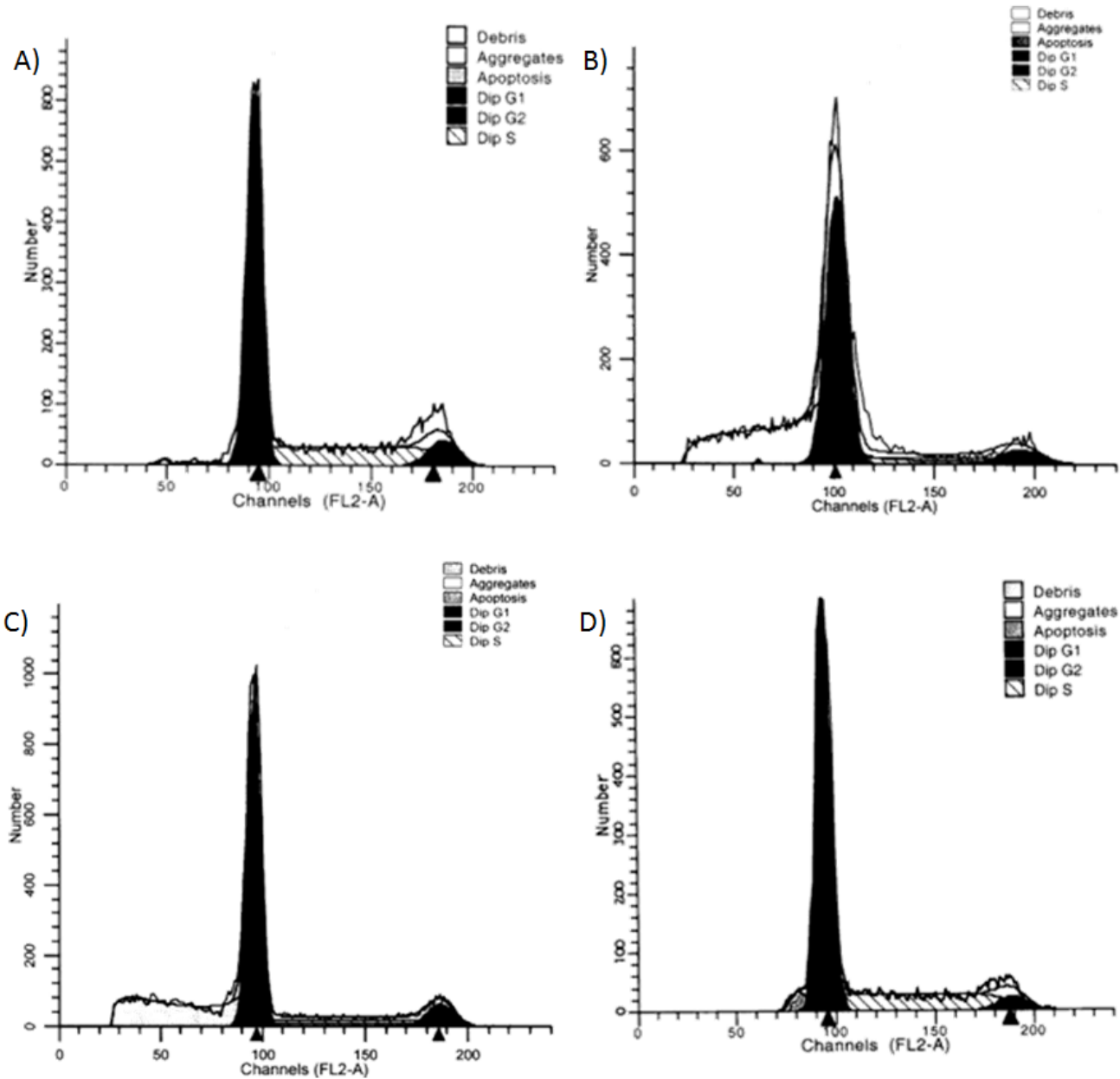


Figure S14. Flow cytometry analysis by propidium iodide of cell cycle arrest in HL60 cells. A) N. C. control; B) cisplatin; C) compound **1** with light irradiation; D) compound **1** in dark. HL60 cells were treated with compounds for 24 hours.

References:

1. Hostetter, A. A.; Osborn, M. F.; DeRose, V. J., RNA-Pt adducts following cisplatin treatment of *Saccharomyces cerevisiae*. *ACS Chem Biol* **2012**, *7* (1), 218-25.
2. Heidary, D. K.; Glazer, E. C., A light-activated metal complex targets both DNA and RNA in a fluorescent in vitro transcription and translation assay. *ChemBiochem* **2014**, *15* (4), 507-11.